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<110> Wood, Clive
Chaudhary, Divya
Long, Andrew

<120> TRADE MOLECULES, AND USES RELATED THERETO

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<150> 60/181,922
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Leu Val Leu Leu Gly Tyr Leu Ser Cys Lys Val Thr Cys Glu Ser Gly
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gac tgt aga cag caa gaa ttc agg gat cgg tct gga aac tgt gtt ccc   144
Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro
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tgc aac cag tgt ggg cca ggc atg gag ttg tct aag gaa tgt ggc ttc   192
Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
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ggc tat ggg gag gat gca cag tgt gtg acg tgc cgg ctg cac agg ttc   240
Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe
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aag gag gac tgg ggc ttc cag aaa tgc aag ccc tgt ctg gac tgc gca   288
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atc tgc ggg gac tgc ttg cca gga ttt tat agg aag acg aaa ctt gtc   384
Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
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ggc ttt caa gac atg gag tgt gtg cct tgt gga gac cct cct cct cct   432
Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
    130                 135                 140

tac gaa ccg cac tgt gcc agc aag gtc aac ctc gtg aag atc gcg tcc   480
Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
145                 150                 155                 160

acg gcc tcc agc cca cgg gac acg gcg ctg gct gcc gtt atc tgc agc   528
Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
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gct ctg gcc acc gtc ctg ctg gcc ctg ctc atc ctc tgt gtc atc tat   576
Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
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tgt aag aga cag ttt atg gag aag aaa ccc agc tgg tct ctg cgg tca   624
Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
        195                 200                 205

cag gac att cag tac aac ggc tct gag ctg tcg tgt ttt gac aga cct   672
Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Phe Asp Arg Pro
    210                 215                 220

cag ctc cac gaa tat gcc cac aga gcc tgc tgc cag tgc cgc cgt gac   720
Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
225                 230                 235                 240

tca gtg cag acc tgc ggg ccg gtg cgc ttg ctc cca tcc atg tgc tgt   768
Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
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gag gag gcc tgc agc ccc aac ccg gcg act ctt ggt tgt ggg gtg cat   816
Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
            260                 265                 270

tct gca gcc agt ctt cag gca aga aac gca ggc cca gcc ggg gag atg   864
Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
        275                 280                 285

gtg ccg act ttc ttc gga tcc ctc acg cag tcc atc tgt ggc gag ttt   912
Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
    290                 295                 300

tca gat gcc tgg cct ctg atg cag aat ccc atg ggt ggt gac aac atc   960
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tct ttt tgt gac tct tat cct gaa ctc act gga gaa gac att cat tct   1008
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caa gat ttg gtt ggt ggg gct gtt cca gtc cag tct cat tct gaa aac   1104
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tca gca tca act cag gat gca cta act atg aga agc cag cta gat cag   1200
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gag agt ggc gct atc atc cac cca gcc act cag acg tcc ctc cag gaa   1248
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gtactccttt gttaggctta tggactgagc agtctggacc ttgcatggct tctggggcaa 1361

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         35                  40                  45

Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
     50                  55                  60
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Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe
 65                  70                  75                  80

Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
                 85                  90                  95

Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
            100                 105                 110

Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
        115                 120                 125

Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
    130                 135                 140

Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
145                 150                 155                 160

Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
                165                 170                 175

Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
            180                 185                 190

Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
        195                 200                 205

Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Phe Asp Arg Pro
    210                 215                 220

Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
225                 230                 235                 240

Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
                245                 250                 255

Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
            260                 265                 270

Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
        275                 280                 285

Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
    290                 295                 300

Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile
305                 310                 315                 320

Ser Phe Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Ile His Ser
                325                 330                 335

Leu Asn Pro Glu Leu Glu Ser Ser Thr Ser Leu Asp Ser Asn Ser Ser
            340                 345                 350

Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn
        355                 360                 365
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Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val Glu
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Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln
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tta gta tta cta ggc tat ttg tca tgt aaa gtg act tgt gaa aca gga   96
Leu Val Leu Leu Gly Tyr Leu Ser Cys Lys Val Thr Cys Glu Thr Gly
            20                  25                  30

gac tgt aga cag caa gaa ttc agg gat cgg tct gga aac tgt gtt ccc   144
Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro
        35                  40                  45

tgc aac cag tgt ggg cca ggc atg gag ttg tct aag gaa tgt ggc ttc   192
Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
    50                  55                  60

ggc tat ggg gag gat gca cag tgt gtg acg tgc cgg ctg cac agg ttc   240
Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe
65                  70                  75                  80

aag gag gac tgg ggc ttc cag aaa tgc aag ccc tgt ctg gac tgc gca   288
Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
                85                  90                  95

gtg gtg aac cgc ttt cag aag gca aat tgt tca gcc acc agt gat gcc   336
Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
            100                 105                 110

atc tgc ggg gac tgc ttg cca gga ttt tat agg aag acg aaa ctt gtc   384
Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
        115                 120                 125

ggc ttt caa gac atg gag tgt gtg cct tgt gga gac cct cct cct cct   432
Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
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    130                 135                 140

tac gaa ccg cac tgt gcc agc aag gtc aac ctc gtg aag atc gcg tcc   480
Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
145                 150                 155                 160

acg gcc tcc agc cca cgg gac acg gcg ctg gct gcc gtt atc tgc agc   528
Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
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gct ctg gcc acc gtc ctg ctg gcc ctg ctc atc ctc tgt gtc atc tat   576
Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
            180                 185                 190

tgt aag aga cag ttt atg gag aag aaa ccc agc tgg tct ctg cgg tca   624
Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
        195                 200                 205

cag gac att cag tac aac ggc tct gag ctg tcg tgt ctt gac aga cct   672
Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Leu Asp Arg Pro
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cag ctc cac gaa tat gcc cac aga gcc tgc tgc cag tgc cgc cgt gac   720
Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
225                 230                 235                 240

tca gtg cag acc tgc ggg ccg gtg cgc ttg ctc cca tcc atg tgc tgt   768
Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
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gag gag gcc tgc agc ccc aac ccg gcg act ctt ggt tgt ggg gtg cat   816
Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
            260                 265                 270

tct gca gcc agt ctt cag gca aga aac gca ggc cca gcc ggg gag atg   864
Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
        275                 280                 285

gtg ccg act ttc ttc gga tcc ctc acg cag tcc atc tgt ggc gag ttt   912
Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
    290                 295                 300

tca gat gcc tgg cct ctg atg cag aat ccc atg ggt ggt gac aac atc   960
Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile
305                 310                 315                 320

tct ttt tgt gac tct tat cct gaa ctc gct gga gaa gac att cat tct   1008
Ser Phe Cys Asp Ser Tyr Pro Glu Leu Ala Gly Glu Asp Ile His Ser
                325                 330                 335

ctc aat cca gaa ctt gaa agc tca acg tct ttg gat tca aat agc agt   1056
Leu Asn Pro Glu Leu Glu Ser Ser Thr Ser Leu Asp Ser Asn Ser Ser
            340                 345                 350

caa gat ttg gtt ggt ggg gct gtt cca gtc cag tct cat tct gaa aac   1104
Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn
        355                 360                 365
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ttt aca gca gct act gat tta tct aga tat aac aac aca ctg gta gaa   1152
Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val Glu
    370                 375                 380

tca gca tca act cag gat gca cta act atg aga agc cag cta gat cag   1200
Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln
385                 390                 395                 400

gag agt ggc gct atc atc cac cca gcc act cag acg tcc ctc cag gta   1248
Glu Ser Gly Ala Ile Ile His Pro Ala Thr Gln Thr Ser Leu Gln Val
                405                 410                 415

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Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro
        35                  40                  45

Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
    50                  55                  60

Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe
65                  70                  75                  80

Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
                85                  90                  95

Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
            100                 105                 110

Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
        115                 120                 125

Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
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Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
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Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
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                165                 170                 175

Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
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Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
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Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Leu Asp Arg Pro
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Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
225                 230                 235                 240

Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
                245                 250                 255

Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
            260                 265                 270

Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
        275                 280                 285

Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
    290                 295                 300

Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile
305                 310                 315                 320

Ser Phe Cys Asp Ser Tyr Pro Glu Leu Ala Gly Glu Asp Ile His Ser
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Leu Asn Pro Glu Leu Glu Ser Ser Thr Ser Leu Asp Ser Asn Ser Ser
            340                 345                 350

Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn
        355                 360                 365

Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val Glu
    370                 375                 380

Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln
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att ctc ttc cta ctc cac ctg gca tgt aaa gtg agt tgc gaa acc gga   96
Ile Leu Phe Leu Leu His Leu Ala Cys Lys Val Ser Cys Glu Thr Gly
             20                  25                  30

gat tgc agg cag cag gaa ttc aag gat cga tct gga aac tgt gtc ctc   144
Asp Cys Arg Gln Gln Glu Phe Lys Asp Arg Ser Gly Asn Cys Val Leu
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tgc aaa cag tgc gga cct ggc atg gag ttg tcc aag gaa tgt ggc ttc   192
Cys Lys Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
     50                  55                  60

ggc tat ggg gag gat gca cag tgt gtg ccc tgc agg ccg cac cgg ttc   240
Gly Tyr Gly Glu Asp Ala Gln Cys Val Pro Cys Arg Pro His Arg Phe
 65                  70                  75                  80

aag gaa gac tgg ggt ttc cag aag tgt aag cca tgt gcg gac tgt gcg   288
Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Ala Asp Cys Ala
                 85                  90                  95

ctg gtg aac cgc ttt cag agg gcc aac tgc tca cac acc agt gat gct   336
Leu Val Asn Arg Phe Gln Arg Ala Asn Cys Ser His Thr Ser Asp Ala
            100                 105                 110

gtc tgc ggg gac tgc ctg cca gga ttt tac cgg aag acc aaa ctg gtt   384
Val Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
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ggt ttt caa gac atg gag tgt gtg ccc tgc gga gac cca cct cct ccc   432
Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
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tac gaa cca cac tgt acc agc aag gtg aac ctt gtg aag atc tcc tcc   480
Tyr Glu Pro His Cys Thr Ser Lys Val Asn Leu Val Lys Ile Ser Ser
145                 150                 155                 160

acc gtc tcc agc cct cgg gac acg gcg ctg gct gcc gtc atc tgc agt   528
Thr Val Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
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gct ctg gcc acg gtg ctg ctc gcc ctg ctc atc ctg tgt gtc atc tac   576
Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
            180                 185                 190

tgc aag agg cag ttc atg gag aag aaa ccc agc tgg tct ctg cgg tca   624
Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
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cag gac att cag tac aat ggc tct gag ctg tca tgc ttt gac cag cct   672
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Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Phe Asp Gln Pro
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cgg ctc cgc cac tgt gcc cat aga gca tgc tgt cag tat cac cgg gac   720
Arg Leu Arg His Cys Ala His Arg Ala Cys Cys Gln Tyr His Arg Asp
225                 230                 235                 240

tca gcc cca atg tat ggg cct gtt cac ctg att ccg tcc ttg tgc tgt   768
Ser Ala Pro Met Tyr Gly Pro Val His Leu Ile Pro Ser Leu Cys Cys
                245                 250                 255

gaa gag gcc cgc agc tct gcc cga gct gtg ctt ggc tgt ggg ctg cgt   816
Glu Glu Ala Arg Ser Ser Ala Arg Ala Val Leu Gly Cys Gly Leu Arg
            260                 265                 270

tct ccc act acc ctc cag gag aga aac ccg gct tct gtg ggg gac acg   864
Ser Pro Thr Thr Leu Gln Glu Arg Asn Pro Ala Ser Val Gly Asp Thr
        275                 280                 285

atg cca gcc ttc ttc ggg tct gtt tcc cgt tcc atc tgc gct gaa ttt   912
Met Pro Ala Phe Phe Gly Ser Val Ser Arg Ser Ile Cys Ala Glu Phe
    290                 295                 300

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Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Leu Gly Gly Asp Ser Ser
305                 310                 315                 320

ctc tgt gac tct tat cct gaa ctc act gga gaa gat acc aat tcc ctc   1008
Leu Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Thr Asn Ser Leu
                325                 330                 335

aat ccc gaa aac gaa agc gca gca tct ctg gat tcc agt ggc ggc cag   1056
Asn Pro Glu Asn Glu Ser Ala Ala Ser Leu Asp Ser Ser Gly Gly Gln
            340                 345                 350

gat ctg gct ggg aca gct gct cta gag tct tct ggg aat gtt tca gaa   1104
Asp Leu Ala Gly Thr Ala Ala Leu Glu Ser Ser Gly Asn Val Ser Glu
        355                 360                 365

tct act gac tca cct aga cat ggt gac act ggt aca gtc tgg gag cag   1152
Ser Thr Asp Ser Pro Arg His Gly Asp Thr Gly Thr Val Trp Glu Gln
    370                 375                 380

acg cta gct cag gat gct caa agg act cca agc caa gga ggc tgg gaa   1200
Thr Leu Ala Gln Asp Ala Gln Arg Thr Pro Ser Gln Gly Gly Trp Glu
385                 390                 395                 400

gac agg gaa aac ctg aat cta gcc atg ccc aca gcc ttc cag gat gcc   1248
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35 40 45

Cys Lys Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
50 55 60

Gly Tyr Gly Glu Asp Ala Gln Cys Val Pro Cys Arg Pro His Arg Phe
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Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Ala Asp Cys Ala
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Leu Val Asn Arg Phe Gln Arg Ala Asn Cys Ser His Thr Ser Asp Ala
100 105 110

Val Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
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Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
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Tyr Glu Pro His Cys Thr Ser Lys Val Asn Leu Val Lys Ile Ser Ser
145 150 155 160

Thr Val Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
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Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
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Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
        195                 200                 205

Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Phe Asp Gln Pro
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Arg Leu Arg His Cys Ala His Arg Ala Cys Cys Gln Tyr His Arg Asp
225                 230                 235                 240

Ser Ala Pro Met Tyr Gly Pro Val His Leu Ile Pro Ser Leu Cys Cys
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Glu Glu Ala Arg Ser Ser Ala Arg Ala Val Leu Gly Cys Gly Leu Arg
            260                 265                 270

Ser Pro Thr Thr Leu Gln Glu Arg Asn Pro Ala Ser Val Gly Asp Thr
        275                 280                 285

Met Pro Ala Phe Phe Gly Ser Val Ser Arg Ser Ile Cys Ala Glu Phe
    290                 295                 300

Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Leu Gly Gly Asp Ser Ser
305                 310                 315                 320

Leu Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Thr Asn Ser Leu
                325                 330                 335

Asn Pro Glu Asn Glu Ser Ala Ala Ser Leu Asp Ser Ser Gly Gly Gln
            340                 345                 350

Asp Leu Ala Gly Thr Ala Ala Leu Glu Ser Ser Gly Asn Val Ser Glu
        355                 360                 365

Ser Thr Asp Ser Pro Arg His Gly Asp Thr Gly Thr Val Trp Glu Gln
    370                 375                 380

Thr Leu Ala Gln Asp Ala Gln Arg Thr Pro Ser Gln Gly Gly Trp Glu
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Asp Arg Glu Asn Leu Asn Leu Ala Met Pro Thr Ala Phe Gln Asp Ala
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